

GBP-1 1 MASEIHM TGPMCLIENTNGRLMANPEALKILSAITQPMVVVAIVGLYRTG
 GBP-2 1 MAPEIINLP GPM SLIDNTKGQLV VNPEALKILSAITQPVVVVAIVGLYRTG
 GBP-4 1 MALEIHM SDPMCLIENT FNEQLKVNQEAL EILSAITQPVVVVAIVGLYRTG

GBP-1 51 KSYLMNKL AGKKKGFS LGSTVQSHTKGIWMWCVPHPKKP GHILVLLDTEG
 GBP-2 51 KSYLMNKL AGKKKGFS LGSTVQSHTKGIWMWCVPHPKKP EHTLVLLDTEG
 GBP-4 51 KSYLMNKL AGKNKGFS VASTVQSHTKGIW IWCVPHPNWPNHTLVLLDTEG

GBP-1 101 LGDVEKGDNDND SWIFALAVLLSSTFVYNS IGTINQQAMDQLYYVTELTH
 GBP-2 101 LGDTEKGDNDND SWIFALAILLSSTFVYNS MGTINQQAMDQLHYVTELTD
 GBP-4 101 LGDVEKADNKNDI QIFALAILLSSTFVYNTVNKIDQGAIDLHNVTELTD

GBP-1 151 RIRSKSSPDENENEVEDSAD FVSFFPDFVWTLRDFS LDLEADGQPLTPDE
 GBP-2 151 RIKANSSPG--NNSVD DSAD FVSFFPAFVWTLRDF TLELEVDGEPI TADD
 GBP-4 151 LLKARNSPD--LDRVED PADSA SFFPD LVWTLRDF CLGLEIDGQLVTPDE

GBP-1 201 YLTYS LKLKKGTSQKDET FNLPRLCIRKFFPKKKCFVFD RPVHRRKLAQL
 GBP-2 199 YLELS LKLKKGTDKKS KSFNDPRLCIRKFFPKR KCFVFDWPAPKKYLAHL
 GBP-4 199 YLENSLRPKQGS DQRVQNFNLPRLCIQKFFPKKKCF TFDLPAHQKKLAQL

GBP-1 251 EKLQDEELDPEFVQQVADFC SYIFSN SKTKTLSGGIQVNGPRLESVLTY
 GBP-2 249 EQLKEEELNPD FIEQVAEFCSYI LSHSNVKTLSGGI AVNGPRLESVLTY
 GBP-4 249 ETLPDDELEPEFVQQVTEFC SYIFSHSM TKTLPGGIMVNGSRLKNLVLTY

GBP-1 301 VNAISSGDLPCMENAVLALAQI ENSAAVQKAI AHYEQQMGQKVQLPTE SL
 GBP-2 299 VNAISSGDLPCMENAVLALAQI ENSAAV EKAIAHYEQQMGQKVQLPTETL
 GBP-4 299 VNAISSGDLPCI TENAVLALAQRENSAAVQKAI AHYDQQMGQKVQLPMETL

GBP-1 351 QELLDLHRDSEREAIEVFIR SSFKDVDH LFOKELAAQLEK KRDDDFCKQNO
 GBP-2 349 QELLDLHRDSEREAIEVF MKNSFKDVDQMFORKLGAQLEAR RDDDFCKONS
 GBP-4 349 QELLDLHRTS EREAIEVF MKNSFKDVDQS FQKELETLLDAKOND ICKRNL

GBP-1 401 EASSDRCSGLLQVIFSPLEEEVKAGIYSKPGGYRLFVQKLQDLKKKYYEE
 GBP-2 399 KASSDCCMALLQDIFGPLEEDVKQGTFSKPGGYRLFTQKLQELKNKYYQV
 GBP-4 399 EASSDYCSALLKDIFGPLEEAVKQGIYSKPGGHNLFIQKTEELKAKYYRE

GBP-1 451 PRKGIQAEELILOTYLSKESMTDAILOTDQTLTEKEKEIEVERVKAESAQ
 GBP-2 449 PRKGIQAEVLK KYLESKEDVADALLQTDQSLSEKEKAIEVERIKAESA E
 GBP-3 1 -----ECVKAESAQ
 GBP-4 449 PRKGIQAEV LQKYLKSKESVSHAILQTDQALTETEKKKKEAQVKAEAEK

GBP-1 501 ASAKMLQEMQRKNEQMMEQKERSYQEH LKQLTEKMENDRVQLLKEQERTL
 GBP-2 499 AAKMLEEIQK KNEEMMEQKEKSYQEHVKQLTEKMERDRAQLMAEQEKTL
 GBP-3 10 ASAKMVEEQIKYQQMMEEKEKSYQEHVKQLTEKMENDRVQLLKEQERTL
 GBP-4 499 AEAQRLAAIQRQNEQMMQERERLHQEQVRQMEIAKQNWLAEQQKMQEQQM

GBP-1 551 ALKLQEQEQLLKEGFQKESRIMKNEIQDLQTKMRRRKA-CTIS
 GBP-2 549 ALKLQEQERLLKEGFENESKRLQKDIWDIQMRSKSLEPICNIL
 GBP-3 60 ALKLQ-----****
 GBP-4 549 QVFINC FISP LPVTMRVCSSGKEGEAARSCGSQQGVWSQKVWV

FIG. 2

mouse.purine.nuc.BP	1	-----MEAPICLVENWKNQLTVNLEAIRILEQIAQPLVVVAIVGL
mu.GBP1	1	--MASEI--HMSEPMCLIENTEAQLVINQEAIRILSAITQPVVVVAIVGL
rat.GBP.p67	1	MDMASEI--HMLQPMCLIENTEAHLVINQEAIRILSAINQPVVVVAIVGL
pig.GBP-1.partial	1	--NFHGIWSTTMDPIXLVKNQNNHLLTVNPKALKILGEICOPVVVVITAGL
pig.GBP-1.NH2.term.	1	--MASKV--HMPEPOCLIENINGRLAVNPKALKILSAIKOPTVVVAIVGL
chicken.GBP	1	--MDTPV-LMPAPLRRLVTNKDGVLAALNTAALAVLRSVTOPVVVAIVAGP
GBP-4	1	--MALEI--HMSDPMCLIENFNEQLKVNQEALEILSAITQPVVVVAIVGL
mouse.purine.nuc.BP	41	YRTGKSYLMNRLAGRNHGFSLGSTVQSETKGIWMWCVPHPTKPTHTLVLL
mu.GBP1	47	YRTGKSYLMNKLAKGRTGFSLGSTVQSHTKGIWMWCVPHPKKAGOTLVLL
rat.GBP.p67	49	YRTGKSYLMNKLAKGRTGFSLGSTVQSHTKGIWMWCVPHPKKAGOTLVLL
pig.GBP-1.partial	49	YRTGKSYLMNRLAGQNHGFRLGSTVRSETKGIWMWCVPHPKKXDHITLVLL
pig.GBP-1.NH2.term.	47	YRTGKSYLMNKLAKKNGK-----
chicken.GBP	48	YRTGKSFLMNRLAQKRTGFPLGPTVYAETKGIWMWCLPHPRQPRVTLVLL
GBP-4	47	YRTGKSYLMNKLAKGKNGFSVASTVQSHTKGIWITWCVPHPNWPNHTLVLL
mouse.purine.nuc.BP	91	DTEGLGDVEKGDPKNDSWIFALAVLLSSTFVYNSMSTINQOALEQLHFVT
mu.GBP1	97	DTEGLEDDVEKGDNDNDWIFALAVLLSSTFYNSIGTINQOAMDQLHYVT
rat.GBP.p67	99	DTEGLEDDVEKGDNDNDWIFALAVLLSSTFVYNSMGITINQOAMDQLHYVT
pig.GBP-1.partial	99	GT-----
chicken.GBP	96	DTEGLEDDPKNDNDHSDAWIFLALLSSTLVYNSVGTIDQALSL---SC
GBP-4	97	DTEGLGDVEKADNKNDIQIFALALLSSTFVYNTVNKIDQGAIDLHNVT
mouse.purine.nuc.BP	141	ELTQLIRAKSSPREDKVKDSSEFVGFFPDFLWAVRDFALELKLNGRPITE
mu.GBP1	147	ELTDLIKSKSSPDQSDVDNSANFVGFFPIFVWTLRDFSLEDFGESITP
rat.GBP.p67	149	ELTDLIKSKSSPDQSGIDDSANFVGFFPTFVWALRDFSLELEVNGKLVTP
chicken.GBP	144	AGNGAVRAHPRGEKDN-NPASNFVSIFPGFVWTVRDFTLQLRDGEKTLSE
GBP-4	147	ELTDLLKARNSPDLDRVEDPADSASFPPDLVWTLRDFCLGLEIDGQLVTP
mouse.purine.nuc.BP	191	DEYLENALKLIQGDNLKVOQSNMTRECIKYFFPVKRCFVFDPRPTSDKRLI
mu.GBP1	197	DEYLETSLALRKGTDENTKKFNMPRLCIRKFFPKRKCFIFDRPGDRKQ-L
rat.GBP.p67	199	DEYLEHSLTLKKGADKKTSGFNEPRLCIRKFFPKRKCFIFDRPALRKQ-L
chicken.GBP	193	DEYLEDVLRIRPGAGRRQERNELRR-CLPNFFPRRKLFITMERPAADAN-L
GBP-4	197	DEYLENSLRPKQSGDQRVQNFNLPRLCIQKFFPKRKCFIFEDLPAHQKK-L
mouse.purine.nuc.BP	241	LQIENVPENQLERNFQVESEKFCSYIFTNGKTKTLRGGVIVITGNRLGTLV
mu.GBP1	246	SKLEWIQEDQLNKEFVEQVAEFTSYIFSYSGVKTLSSGGITVNGPRLKSLV
rat.GBP.p67	248	CKLETTLGEEELCSEFVEQVAEFTSYIFSYSAVKTLSSGGIIVNGPRLKSLV
chicken.GBP	241	TRLEELREDELQPGFRKQVDAFCRYIWEAPVKVLPGGHQVITGSALAYLV
GBP-4	246	AQLETLPDDELPEFVEQVTEFCSYIFSHSMTKTLPGGIMVNGSRLKNLV
mouse.purine.nuc.BP	291	QTYVNAIINSGTVPCLENAVITTLAQRENSIAVOKAADHYSEQMAQRMRLPT
mu.GBP1	296	QTYVSAICSGELPCMENAVLTLAQIENSAAVOKAITYYEEQMNQKIHMPT
rat.GBP.p67	298	QTYVGAISSGSLPCMESAVALTLAQIENSAAVOKAITHYEEQMNQKIOMPT
chicken.GBP	291	EKYMAAISSGSVPCVESTLKALAQAEHTAAVQMAVAEYQRMEOGLVLPT
GBP-4	296	LTYVNAISSGDLPCIENAVLALAQRENSAAVOKAIAHYDOQMCKVQLPM
mouse.purine.nuc.BP	341	DTLQELLTVHAACEKEAIAVFMHSFKDDEQEFQKKLVVTIEERKEEFIR
mu.GBP1	346	ETLQELLDLHRTCEREAEVFMKNSFKDVDQKFQEELEGAQLEAKRDAFVK
rat.GBP.p67	348	ETLQELLDLHRLIEREAIEIFLKNSFKDVDQKFQETELGNLLISKRDAFIK
chicken.GBP	341	ASYDALAVHRDCEQALALFLSRAFAADHKHGYHDELVHKLEGQRGVLP
GBP-4	346	ETLQELLDLHRTSEREAIEVFMKNSFKDVDQSFQKELETLLDAKQNDICK
mouse.purine.nuc.BP	391	QNEAASIRHCQAELERLSESLRKSISCGAFSVPGGHSLYLEARKKIELGY
mu.GBP1	396	KNMDMSSAHCSDLLEGLFAHLEEEVKQGTIFYKPGGYLFLQKQOELEKKY
rat.GBP.p67	398	KNSDVSSAHCSDLIEDIFGPLEEEVKQGTFSKPGGYFLFLQMRQOELEKKY
chicken.GBP	391	QGGVGAAVPHGAAGAVEGR-GRRLQRCGYVARGGAAQLFKEDVNRVLEEY
GBP-4	396	RNLEASSDYCSALLKDDFGPDEEAVKQGIYSKPGGHNLFIQKTEELKAKY

FIG. 3A

mouse.purine.nuc.BP	441	QQVLRKGLKAKEVLKSFLOSQAIMEDSILQSDKALTDGERAIAAERTKKE
mu.GBP1	446	IQTPGKGLQAEVMLRKYFESKEDLADTLKMDQSLTEKEKQIEMERIKAE
rat.GBP.p67	448	NOAPGKGLEAEAVLKKYFESKEDIVETLLKTDQSLTEAAKEIEVERIKAE
pig.GBP-1.partial	101	-----PRG-----
chicken.GBP	440	KQRPDKGVRAEAVLKEFLREHEGLAQVLKATEVOLELAERQOEAAAAEAE
GBP-4	446	YREPRKGIQAEVLOKYLKSKESVSHAILQTDQALTETEKKKKKEAQVKAE
mouse.purine.nuc.BP	491	VAEKELELLRROKKEQEQVMEAQERSFRENIAKLQEKMES---EKEMLLR
mu.GBP1	496	AAEAANRALAEMOKKHEMLMEQKEQSYQEHMKQLTEKMEQ---ERKELMA
rat.GBP.p67	498	TAEAAANRELAEKQEKFLMLMQOKEESYQEHVROLTEKMKE---EQKKLIE
chicken.GBP	490	AARKATEAWREDQK-----RSMEEHKRQLEQWMKK---EKHTWEE
GBP-4	496	AEKAEAQRLAAIQRONEQMMQERERLHQEQVROMETAKQNWLAEEQQKMQE
mouse.purine.nuc.BP	538	EEKMLEHKLKVQEELLIEGFREKSDMLKNETSHLREEMERTRRKPSLFG
mu.GBP1	543	EQQRIISLKLQEQERILLKQGFQNESLQLRQETIEKIK-NMPPPRS-----
rat.GBP.p67	545	EQDNIIALKLREQEKFLREGYENESKKLLREIENMK-RRQSPGK-----
chicken.GBP	527	ELNRMLEHHRKEYKALLQEGFRREAAAKEKQIRELQEEMRSCN-----
GBP-4	546	QQMQVFINC FISPLPVTMRVCSGKEGEAARSCGSQ-QGVWSQK-----
mouse.purine.nuc.BP	588	QILDTIGNAFIMILPGAGKLFGVGLKFLGSLSS
mu.GBP1	586	-----CTIL-----
rat.GBP.p67	588	-----CTIL-----
chicken.GBP	570	-----CTVL-----
GBP-4	589	-----VWV-----

FIG. 3B

hGBP-4						
67	hGBP-1					
64	76	hGBP-2				
60	68	71	rGBP			
59	70	71	80	mGBP-1		
49	54	54	52	53	mPNBP	
37	41	41	41	39	41	cGBP

FIG. 4



FIG. 5

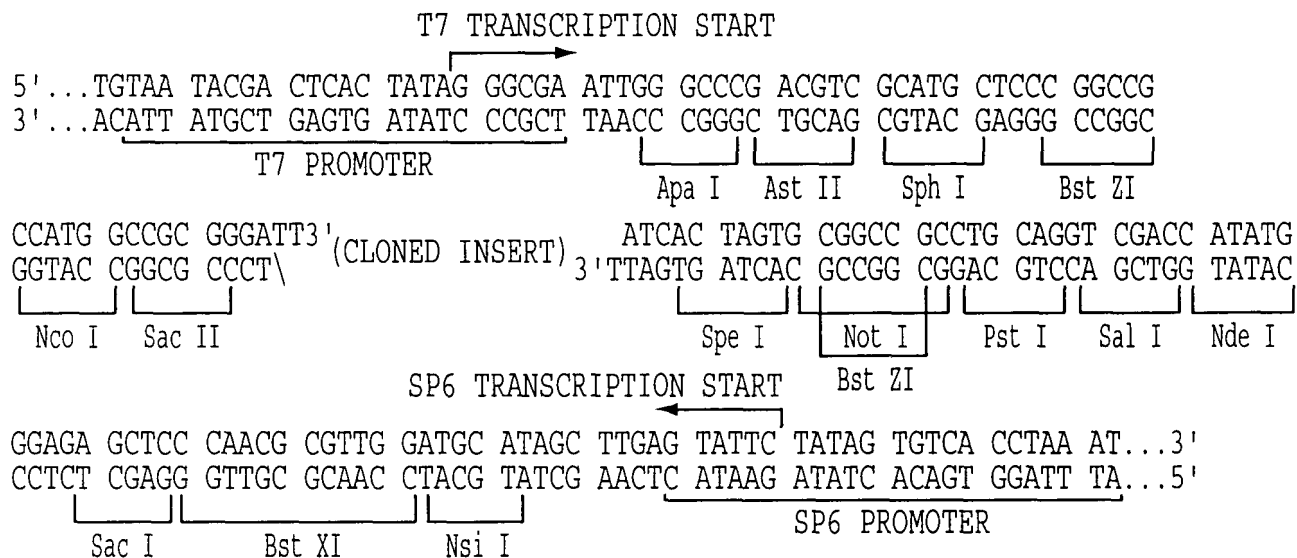


FIG. 6

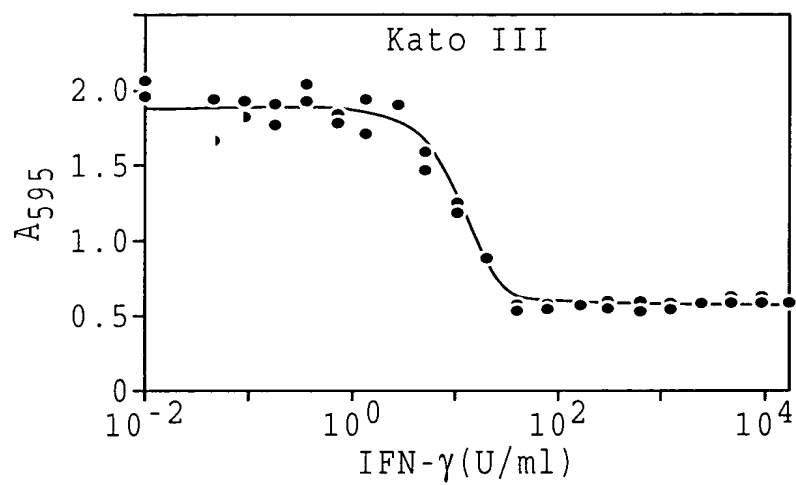


FIG. 7A

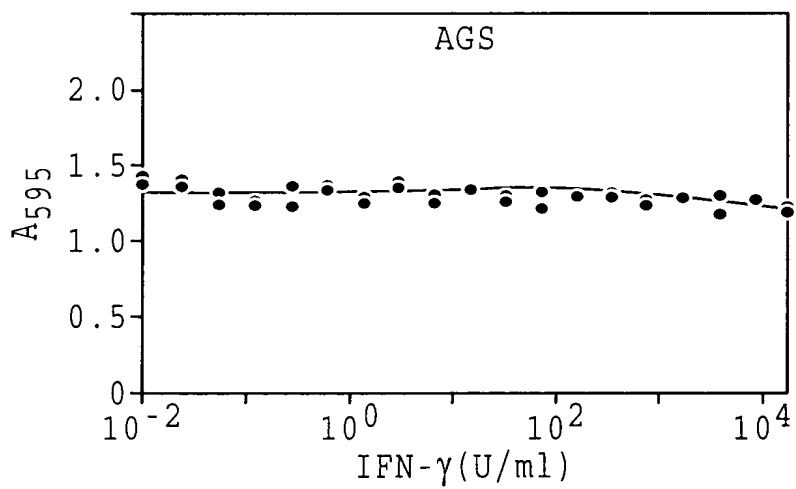


FIG. 7B

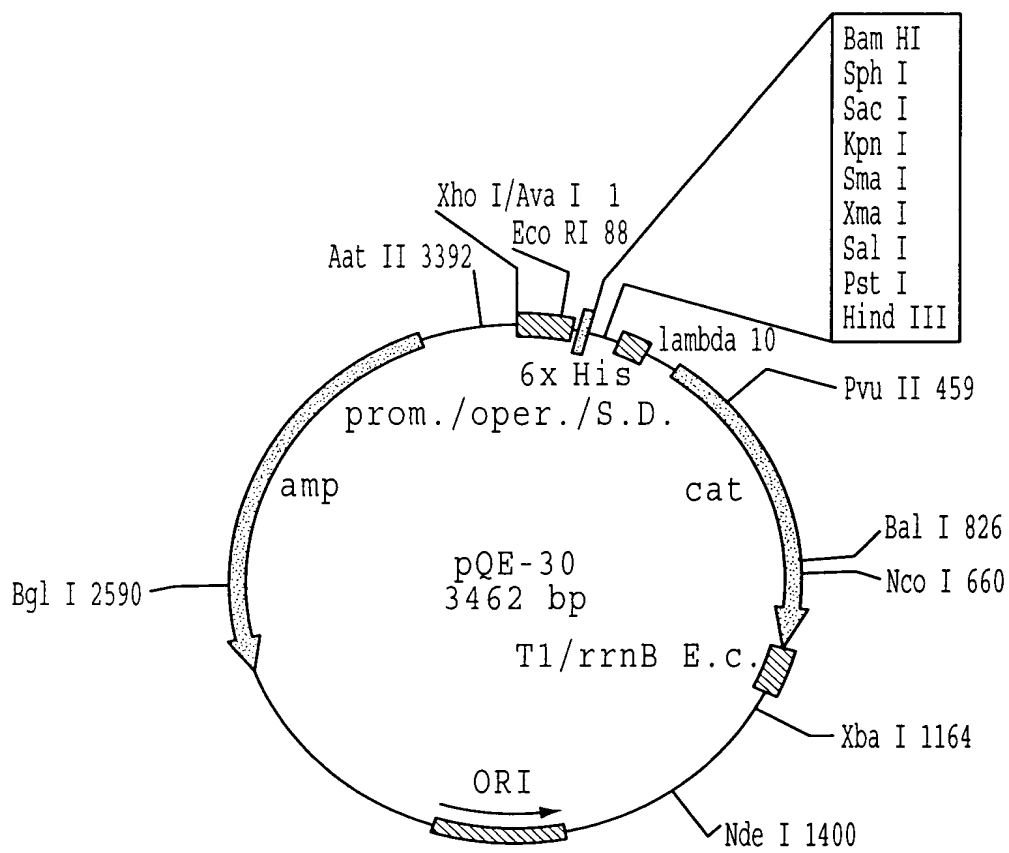


FIG. 8